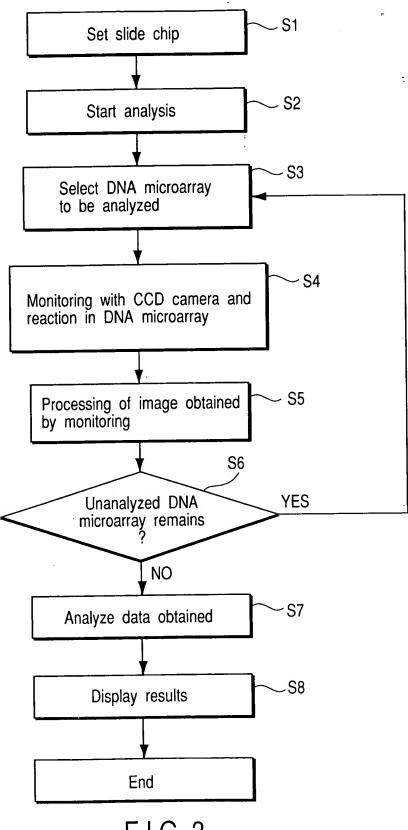
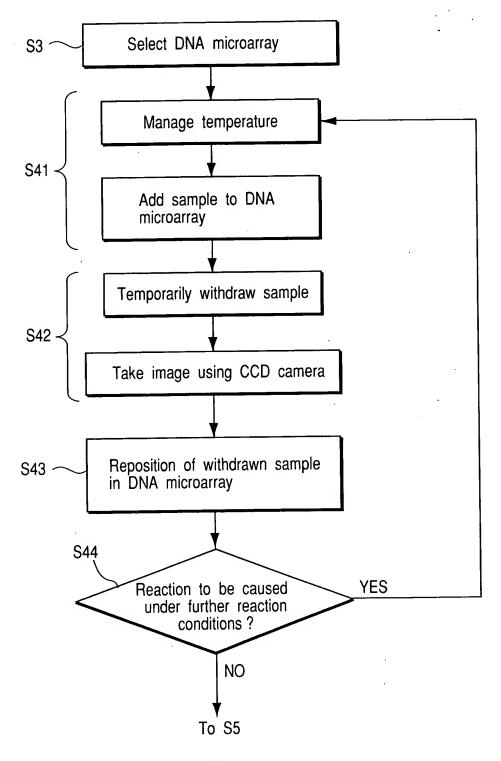


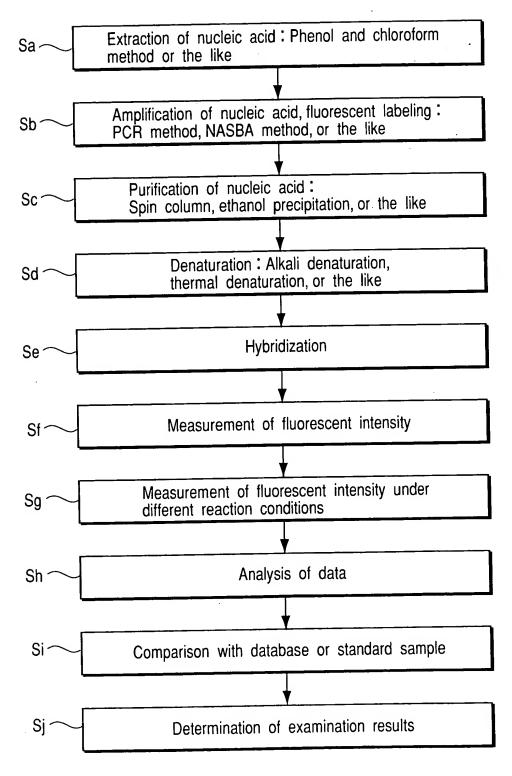
F I G. 2



F I G. 3



F I G. 4

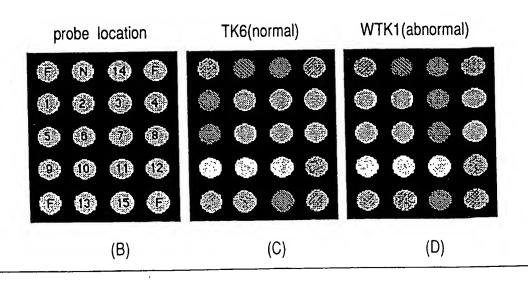


F I G. 5

umber	spot oligo (	25 mer)	target
1	5'-AGTTTGTGTTTCAACTGTTCTCGTC-3'		c-myc
2	ATCTGTCTCAGGACTCTGACACTGT		c-myc
3	ACTCAAACGTGTCTGTGTTGTAGGT		ERBB2
4	AATCTGCATACACCAGTTCAGCAG (24mer)		ERBB2
5	CATAATGGTAGCCTGAAGCATAGTC		ER
	GGATCAAAGTGTCTGTGATCTTGTC		ER
6	TACAGATGAGGTT	TTTGCCTGAGT	ZABC1(ZNF217)
7	ATAACTCTTGATAT	GACACAGGCCT	ZABC1(ZNF218)
.8	ATAAGTGTTGATATGACACAGGCCT CTCGTCTTCTACAGGGAAGTTCAC (24mer)		hTERT
9	CAGGAGGATCTTGTAC	ATGTTGGT (24mer)	hTERT
10			Luciferase gene of Renia reniformis
11	ACATCTACTACAC <sup>*</sup>	TTCAGCGTGAA	(negative control)
		•	Luciferase gene of Renia reniformis
12	CGTCAGGTTTACC	ACCTTTTACTAA	(negative control)
	•		β –actin (positive control)
13	GTCACACTTCATG	TOOTTAATOTOA	β -actin (positive control)
14	GTAGCACAGCTTC	TOTOATACTTCT	GAPDH (positive control)
15	ATCTTGAGGCTGTTGTCATACTTCT		GAPDH (positive control)
16	ACCACCTTCTTGATGTCATCATATT		GAPDH Costave contactive
F	fluorescein oligo		
N	non D	(A)	
	probe location	TIG-1(G)	TIG-1(G)
	0000		
	0 0 0 0		
	3 3 4 4		
	5 5 6 6	0000	
	5 5 6 6 7 7 8 8		
	7 7 8 8 9 9 10 0		
	0 0 8 8		
	7 7 8 8 9 9 10 0 11 (1) 12 12		
	7 7 8 8 9 9 10 0		
	7 7 8 8 9 9 10 0 11 11 12 12 13 13 14 14		
	7 7 8 8 9 9 10 0 11 (1) 12 12		
	7 7 8 8 9 9 10 0 11 11 12 12 13 13 14 14 15 15 16 16		
	7 7 8 8 9 9 10 0 11 11 12 12 13 13 14 14		
	7 7 8 8 9 9 10 0 11 11 12 12 13 13 14 14 15 15 16 16	(C)	

number	spot oligo (21 mer)	number of mismatch
1	5'-ACAACTACATGTGTAACAGTT-3'	0 (sense)
2	ACAACTACATCTGTAACAGTT	1
3	ACAACTACATATGTAACAGTT	1
4	ACAACTACATTTGTAACAGTT	1
5	AACTGTTACACATGTAGTTGT	0 (antisense)
6	AACTGTTACAGATGTAGTTGT	1
. 7	AACTGTTACATATGTAGTTGT	1
8	AACTGTTACAAATGTAGTTGT	1
9	ACAACTACAGATGTAACAGTT	2
10	ACAACTACATATGTAGCAGTT	, <b>2</b>
11	ACAAGTACATATGTAACAGTT	2
12	ACAAGTACATATGTAGCAGTT	3_
13	ACAAGTACAGACGTAGCAGTT	5
14	CACAGGCCCAAGATGAGGCC	complement of primer F
15	ACTTGCCACCCTGCACACTG	complement of primer R
F	fluorescein oligo	
N	non DNA	

(A)



F I G. 7

